RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: _10/809,945_

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RAW SEQUENCE LISTING   DATE: 12/01/2004
PATENT APPLICATION: **US/10/809,945**   TIME: 11:56:32

Input Set : **N:\Crf3\RULE60\10809945.raw.txt**
Output Set: **N:\CRF4\12012004\J809945.raw**

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 1 <110> APPLICANT: Harberd, Nicholas P
 2         Richards, Donald E
 3         Peng, Jinrong
 4 <120> TITLE OF INVENTION: Genetic Control of Plant Growth and Development
 5 <130> FILE REFERENCE: 620-91
 6 <140> CURRENT APPLICATION NUMBER: US/10/809,945
 7 <141> CURRENT FILING DATE: 2004-03-26
 8 <150> PRIOR APPLICATION NUMBER: US/09/485,529
 9 <151> PRIOR FILING DATE: 2000-03-01
10 <150> PRIOR APPLICATION NUMBER: PCT/GB98/02383
11 <151> PRIOR FILING DATE: 1998-08-07
12 <150> PRIOR APPLICATION NUMBER: GB 9717192.0
13 <151> PRIOR FILING DATE: 1997-08-13
14 <160> NUMBER OF SEQ ID NOS: 108
15 <170> SOFTWARE: PatentIn Ver. 2.0
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 630
19 <212> TYPE: PRT
20 <213> ORGANISM: Triticum aestivum
21 <220> FEATURE:
22 <221> NAME/KEY: SITE
23 <222> LOCATION: (91)
24 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
25 <221> NAME/KEY: SITE
26 <222> LOCATION: (94)
27 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
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29 <222> LOCATION: (100)
30 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
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32 <222> LOCATION: (106)
33 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
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35 <222> LOCATION: (118)
36 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
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38 <222> LOCATION: (121)
39 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W--> 40 <221> SITE
41 <222> LOCATION: (142)
42 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W--> 43 <221> SITE
44 <222> LOCATION: (197)
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        47 <222> LOCATION: (436)
        48 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    49 <221> SITE
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        51 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    52 <221> SITE
        53 <222> LOCATION: (474)
        54 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    55 <221> SITE
        56 <222> LOCATION: (536)
        57 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    58 <221> SITE
        59 <222> LOCATION: (558)
        60 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    61 <221> SITE
        62 <222> LOCATION: (589)
        63 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    64 <221> SITE
        65 <222> LOCATION: (602)
        66 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    67 <221> SITE
        68 <222> LOCATION: (617)
        69 <223> OTHER INFORMATION: Xaa is unknown or other amino acid
W-->    70 <400> 1
        71      Ile Glu Arg Arg Gly Ser Ser Arg Ile Met Lys Arg Glu Tyr Gln Asp
        72      1               5                   10                  15
        73      Ala Gly Gly Ser Gly Gly Gly Gly Gly Gly Met Gly Ser Glu Asp Lys
        74                      20                  25              30
        75      Met Met Val Ser Ala Ala Ala Gly Glu Gly Glu Glu Val Asp Glu Leu
        76                  35              40                  45
        77      Leu Ala Ala Leu Gly Tyr Lys Val Arg Ala Ser Asp Met Ala Asp Val
        78          50                  55                  60
        79      Ala Gln Lys Leu Glu Lys Leu Glu Met Ala Met Gly Met Gly Gly Val
        80          65                  70                  75                  80
W-->    81      Gly Ala Gly Ala Ala Pro Asp Arg Gln Val Xaa His Pro Xaa Ala Ala
        82                      85                  90                  95
        83      Asp Thr Val Xaa Tyr Asn Pro Thr Asp Xaa Ser Ser Trp Val Glu Ser
        84                      100                 105                 110
        85      Met Leu Ser Glu Leu Xaa Glu Pro Xaa Pro Pro Leu Pro Pro Ala Pro
        86                  115                 120                 125
        87      Gln Leu Asn Ala Ser Thr Val Thr Gly Ser Gly Gly Tyr Xaa Asp Leu
        88          130                 135                 140
        89      Pro Pro Ser Val Asp Ser Ser Ser Ser Ile Tyr Ala Leu Arg Pro Ile
        90      145                 150                 155                 160
        91      Pro Ser Pro Ala Gly Ala Thr Ala Pro Ala Asp Leu Ser Ala Asp Ser
        92                      165                 170                 175
        93      Val Arg Asp Pro Lys Arg Met Arg Thr Gly Gly Ser Ser Thr Ser Ser
```

```
94                  180                   185                   190
95    Ser Ser Ser Ser Xaa Ser Ser Leu Gly Gly Gly Ala Arg Ser Ser Val
96              195                   200                   205
97    Val Glu Ala Ala Pro Pro Val Ala Ala Ala Ala Asn Ala Thr Pro Ala
98        210                   215                   220
99    Leu Pro Val Val Val Val Asp Thr Gln Glu Ala Gly Ile Arg Leu Val
100   225                   230                   235                   240
101     His Ala Leu Leu Ala Cys Ala Glu Ala Val Gln Gln Glu Asn Leu Ser
102               245                   250                   255
103     Ala Ala Glu Ala Leu Val Lys Gln Ile Pro Leu Leu Ala Ala Ser Gln
104               260                   265                   270
105     Gly Gly Ala Met Arg Lys Val Ala Ala Tyr Phe Gly Glu Ala Leu Ala
106               275                   280                   285
107     Arg Arg Val Phe Arg Phe Arg Pro Gln Pro Asp Ser Ser Leu Leu Asp
108               290                   295                   300
109     Ala Ala Phe Ala Asp Leu Leu His Ala His Phe Tyr Glu Ser Cys Pro
110   305                   310                   315                   320
111   Tyr Leu Lys Phe Ala His Phe Thr Ala Asn Gln Ala Ile Leu Glu Ala
112               325                   330                   335
113   Phe Ala Gly Cys Arg Arg Val His Val Val Asp Phe Gly Ile Lys Gln
114               340                   345                   350
115   Gly Met Gln Trp Pro Ala Leu Leu Gln Ala Leu Ala Leu Arg Pro Gly
116               355                   360                   365
117   Gly Pro Pro Ser Phe Arg Leu Thr Gly Val Gly Pro Pro Gln Pro Asp
118               370                   375                   380
119   Glu Thr Asp Ala Leu Gln Gln Val Gly Trp Lys Leu Ala Gln Phe Ala
120   385                   390                   395                   400
121   His Thr Ile Arg Val Asp Phe Gln Tyr Arg Gly Leu Val Ala Ala Thr
122               405                   410                   415
123   Leu Ala Asp Leu Glu Pro Phe Met Leu Gln Pro Glu Gly Glu Glu Asp
124               420                   425                   430
125   Pro Asn Glu Xaa Pro Xaa Val Ile Ala Val Asn Ser Val Phe Glu Met
126               435                   440                   445
127   His Arg Leu Leu Ala Gln Pro Gly Ala Leu Glu Lys Val Leu Gly His
128         450                   455                   460
129   Arg Ala Pro Pro Cys Gly Pro Glu Phe Xaa Thr Val Val Glu Thr Gln
130   465                   470                   475                   480
131   Glu Ala Asn His Asn Ser Gly Thr Phe Leu Asp Arg Phe Thr Glu Ser
132               485                   490                   495
133   Leu His Tyr Tyr Ser Thr Met Phe Asp Ser Leu Glu Gly Gly Ser Ser
134               500                   505                   510
135   Gly Gly Gly Pro Ser Glu Val Ser Ser Gly Ala Ala Ala Ala Pro Ala
136               515                   520                   525
137   Ala Ala Gly Thr Asp Gln Val Xaa Ser Glu Val Tyr Leu Gly Arg Gln
138         530                   535                   540
139   Ile Cys Asn Val Val Ala Cys Glu Gly Ala Glu Arg Thr Xaa Arg His
140   545                   550                   555                   560
141   Glu Thr Leu Gly Gln Trp Arg Asn Arg Leu Gly Asn Ala Gly Phe Glu
142               565                   570                   575
```

```
143      Thr Val His Leu Gly Ser Asn Ala Tyr Lys Gln Ala Xaa Thr Leu Leu
144              580                 585                 590
145      Ala Leu Phe Ala Gly Gly Glu Arg Leu Xaa Val Glu Glu Lys Glu Gly
146              595                 600                 605
147      Cys Leu Thr Leu Gly Leu His Thr Xaa Pro Leu Ile Ala Thr Ser Ala
148          610                 615                 620
149      Trp Arg Leu Ala Gly Pro
150      625                 630
152 <210> SEQ ID NO: 2
153 <211> LENGTH: 532
154 <212> TYPE: PRT
155 <213> ORGANISM: Arabidopsis thaliana
156 <400> SEQUENCE: 2
157      Met Lys Arg Asp His His His His His Gln Asp Lys Lys Thr Met Met
158        1                 5                 10                  15
159      Met Asn Glu Glu Asp Asp Gly Asn Gly Met Asp Glu Leu Leu Ala Val
160              20                 25                  30
161      Leu Gly Tyr Lys Val Arg Ser Ser Glu Met Ala Asp Val Ala Gln Lys
162              35                 40                  45
163      Leu Glu Gln Leu Glu Val Met Met Ser Asn Val Gln Glu Asp Asp Leu
164          50                 55                  60
165      Ser Gln Leu Ala Thr Glu Thr Val His Tyr Asn Pro Ala Glu Leu Tyr
166      65                 70                  75                  80
167      Thr Trp Leu Asp Ser Met Leu Thr Asp Leu Asn Pro Pro Ser Ser Asn
168              85                 90                  95
169      Ala Glu Tyr Asp Leu Lys Ala Ile Pro Gly Asp Ala Ile Leu Asn Gln
170              100                105                 110
171      Phe Ala Ile Asp Ser Ala Ser Ser Ser Asn Gln Gly Gly Gly Gly Asp
172              115                120                 125
173      Thr Tyr Thr Thr Asn Lys Arg Leu Lys Cys Ser Asn Gly Val Val Glu
174          130                135                 140
175      Thr Thr Thr Ala Thr Ala Glu Ser Thr Arg His Val Val Leu Val Asp
176      145                 150                 155                 160
177      Ser Gln Glu Asn Gly Val Arg Leu Val His Ala Leu Leu Ala Cys Ala
178              165                170                 175
179      Glu Ala Val Gln Lys Glu Asn Leu Thr Val Ala Glu Ala Leu Val Lys
180              180                185                 190
181      Gln Ile Gly Phe Leu Ala Val Ser Gln Ile Gly Ala Met Arg Lys Val
182              195                200                 205
183      Ala Thr Tyr Phe Ala Glu Ala Leu Ala Arg Arg Ile Tyr Arg Leu Ser
184          210                215                 220
185      Pro Ser Gln Ser Pro Ile Asp His Ser Leu Ser Asp Thr Leu Gln Met
186      225                 230                 235                 240
187      His Phe Tyr Glu Thr Cys Pro Tyr Leu Lys Phe Ala His Phe Thr Ala
188              245                250                 255
189      Asn Gln Ala Ile Leu Glu Ala Phe Gln Gly Lys Lys Arg Val His Val
190              260                265                 270
191      Ile Asp Phe Ser Met Ser Gln Gly Leu Gln Trp Pro Ala Leu Met Gln
192          275                280                 285
```

```
193         Ala Leu Ala Leu Arg Pro Gly Gly Pro Pro Val Phe Arg Leu Thr Gly
194             290                 295                 300
195         Ile Gly Pro Pro Ala Pro Asp Asn Phe Asp Tyr Leu His Glu Val Gly
196         305                 310                 315                 320
197         Cys Lys Leu Ala His Leu Ala Glu Ala Ile His Val Glu Phe Glu Tyr
198                         325                 330                 335
199         Arg Gly Phe Val Ala Asn Thr Leu Ala Asp Leu Asp Ala Ser Met Leu
200                     340                 345                 350
201         Glu Leu Arg Pro Ser Glu Ile Glu Ser Val Ala Val Asn Ser Val Phe
202                     355                 360                 365
203         Glu Leu His Lys Leu Leu Gly Arg Pro Gly Ala Ile Asp Lys Val Leu
204             370                 375                 380
205         Gly Val Val Asn Gln Ile Lys Pro Glu Ile Phe Thr Val Val Glu Gln
206         385                 390                 395                 400
207         Glu Ser Asn His Asn Ser Pro Ile Phe Leu Asp Arg Phe Thr Glu Ser
208                         405                 410                 415
209         Leu His Tyr Tyr Ser Thr Leu Phe Asp Ser Leu Glu Gly Val Pro Ser
210                         420                 425                 430
211         Gly Gln Asp Lys Val Met Ser Glu Val Tyr Leu Gly Lys Gln Ile Cys
212             435                 440                 445
213         Asn Val Val Ala Cys Asp Gly Pro Asp Arg Val Glu Arg His Glu Thr
214             450                 455                 460
215         Leu Ser Gln Trp Arg Asn Arg Phe Gly Ser Ala Gly Phe Ala Ala Ala
216         465                 470                 475                 480
217         His Ile Gly Ser Asn Ala Phe Lys Gln Ala Ser Met Leu Leu Ala Leu
218                         485                 490                 495
219         Phe Asn Gly Gly Glu Gly Tyr Arg Val Glu Glu Ser Asp Gly Cys Leu
220                     500                 505                 510
221         Met Leu Gly Trp His Thr Arg Pro Leu Ile Ala Thr Ser Ala Trp Lys
222                     515                 520                 525
223         Leu Ser Thr Asn
224             530
226 <210> SEQ ID NO: 3
227 <211> LENGTH: 2709
228 <212> TYPE: DNA
229 <213> ORGANISM: Triticum aestivum
230 <220> FEATURE:
231 <221> NAME/KEY: misc_feature
232 <222> LOCATION: (6)
233 <223> OTHER INFORMATION: n is any nucleotide
234 <221> NAME/KEY: misc_feature
235 <222> LOCATION: (11)
236 <223> OTHER INFORMATION: n is any nucleotide
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238 <222> LOCATION: (46)
239 <223> OTHER INFORMATION: n is any nucleotide
W--> 240 <221> misc_feature
241 <222> LOCATION: (85)
242 <223> OTHER INFORMATION: n is any nucleotide
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Input Set : **N:\Crf3\RULE60\10809945.raw.txt**
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:1; Xaa Pos. 91,94,100,106,118,121,142,197,436,438,474,536,558,589,602
Seq#:1; Xaa Pos. 617
Seq#:3; N Pos. 6,11,46,85,107,407,417,430,436,448,452,490,498,556,562,727
Seq#:3; N Pos. 769,1444,1450,1556,1742,1808,1900,1940,1982,2438,2604,2638
Seq#:3; N Pos. 2681,2689,2706
Seq#:19; N Pos. 171,302,427,444,459,711
Seq#:20; Xaa Pos. 57,143,148,250
Seq#:57; N Pos. 674,680,786,972,1038,1130,1170,1212
Seq#:58; N Pos. 19,33,131
Seq#:59; N Pos. 17,19,20,53,61,77,84,137,142,156,162,167,205,212,221,251
Seq#:59; N Pos. 259,271,273,277,290,306,321,337,342,345,362,367,369
Seq#:60; N Pos. 3,34,45,142,176,179,185,186,187,188,189,191,197,198,210
Seq#:61; N Pos. 5,8,11,51,56,59,69,156,210,264,345,350,360,367
Seq#:62; N Pos. 6,57,177,235,238,251,286,293,294,295,296,328,331,332,333
Seq#:63; N Pos. 7,17,149,263,361,362,413,438,479,484,491,496,510
Seq#:64; N Pos. 9,46,50,72,95,131,141,154,160,172,176,214,222
Seq#:65; N Pos. 7,13,17,24,37,45,46,55,60,61,87,108,123,130,135,163,176,233
Seq#:65; N Pos. 347,350,370,378,379,383,386,399
Seq#:66; N Pos. 6,11,46,85,107,445,449
Seq#:67; N Pos. 57,66,127,131,147,201,367,401,444,452,469
Seq#:68; N Pos. 4,9,12,28,31,36,41,42,46,54,56,60,63,92,157,161,173,184,195
Seq#:68; N Pos. 198,200,205,213
Seq#:69; N Pos. 37,66,164,172,179,185,188
Seq#:70; N Pos. 4,14,202,210,214
Seq#:71; N Pos. 2,3,8,10,11,12,13,25,33,39,365
Seq#:72; N Pos. 1,21,54,60,225,267,364,385,394,404,412,415,416,425,426,427
Seq#:72; N Pos. 433,436
Seq#:73; N Pos. 3,11,196,274,369,370,371,379,394,402,414,423
Seq#:74; N Pos. 24,90,121,170,173,194,203,229,255,263,273
Seq#:75; N Pos. 10,13,17,18,45,48,158,169,205,218,224,231,237,242,258
Seq#:76; N Pos. 158,161,217
Seq#:77; N Pos. 38,108,128,352,353,371,383,385
Seq#:79; Xaa Pos. 26,31
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Input Set : **N:\Crf3\RULE60\10809945.raw.txt**
Output Set: **N:\CRF4\12012004\J809945.raw**

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L:28 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:31 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:34 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:37 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:40 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:43 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:46 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:49 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:52 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:58 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:61 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:64 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:67 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:70 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:80
M:341 Repeated in SeqNo=1
L:237 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:240 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:243 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:246 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:249 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:252 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:255 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:258 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:261 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:264 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:267 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:270 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
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L:285 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:288 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
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L:294 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:297 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:300 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
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L:321 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:324 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
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```
M:341 Repeated in SeqNo=3
L:904 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:907 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:910 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:913 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:916 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:919 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:120
M:341 Repeated in SeqNo=19
L:955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:48
M:341 Repeated in SeqNo=20
L:1349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 after pos.:660
M:341 Repeated in SeqNo=57
L:1384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:0
M:341 Repeated in SeqNo=58
L:1481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
M:341 Repeated in SeqNo=59
L:1540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:0
M:341 Repeated in SeqNo=60
L:1593 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0
M:341 Repeated in SeqNo=61
L:1652 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0
M:341 Repeated in SeqNo=62
L:1704 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:0
M:341 Repeated in SeqNo=63
L:1759 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0
M:341 Repeated in SeqNo=64
L:1850 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:0
M:341 Repeated in SeqNo=65
L:1885 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:0
M:341 Repeated in SeqNo=66
L:1933 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:0
M:341 Repeated in SeqNo=67
L:2017 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:0
M:341 Repeated in SeqNo=68
L:2049 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:0
M:341 Repeated in SeqNo=69
L:2075 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0
M:341 Repeated in SeqNo=70
L:2119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71 after pos.:0
M:341 Repeated in SeqNo=71
L:2187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72 after pos.:0
M:341 Repeated in SeqNo=72
L:2238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:0
M:341 Repeated in SeqNo=73
L:2286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0
M:341 Repeated in SeqNo=74
L:2343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75 after pos.:0
M:341 Repeated in SeqNo=75
L:2366 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76 after pos.:120
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VERIFICATION SUMMARY   DATE: 12/01/2004
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   Input Set : **N:\Crf3\RULE60\10809945.raw.txt**
   Output Set: **N:\CRF4\12012004\J809945.raw**

M:341 Repeated in SeqNo=76
L:2401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77 after pos.:0
M:341 Repeated in SeqNo=77